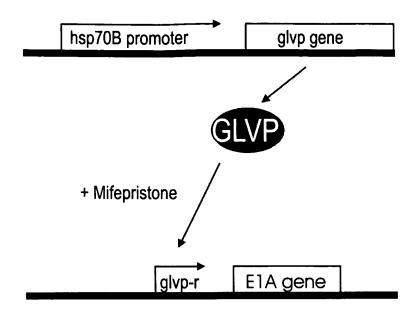
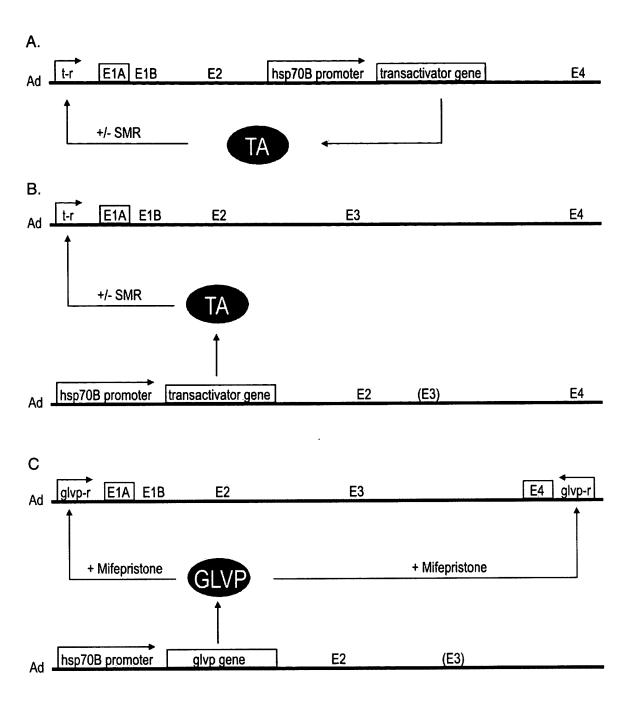


TA - transactivator protein
 t-r - transactivator-responsive promoter
 SMR - small molecule regulator



glvp-r - GLVP-responsive promoter



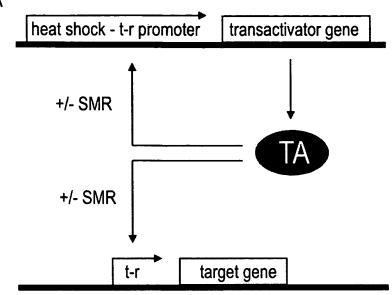
glvp-r - GLVP-responsive promoter

t-r - transactivator-responsive promoter

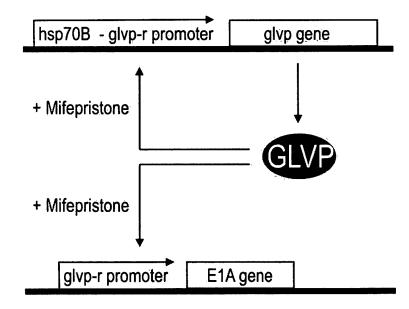
SMR - small molecule regulator TA - transactivator protein

Ad - adenovirus DNA

() - gene region from which sequences can be optionally deleted



В



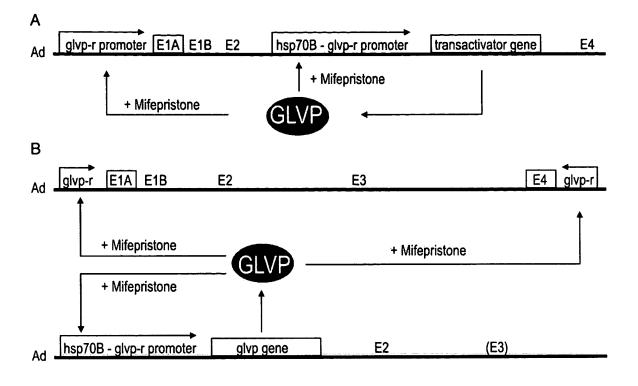
glvp-r - GLVP-responsive promoter

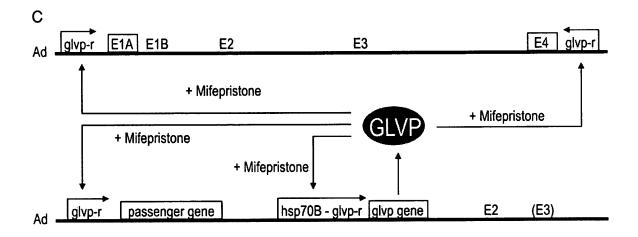
- transactivator-responsive promoter

SMR - small molecule regulator - transactivator protein

hsp70B - glvp-r promoter heat shock - t-r promoter

tandem or hybrid promoters





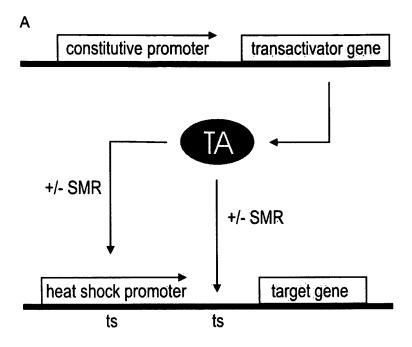
glvp-r - GLVP-responsive promoter

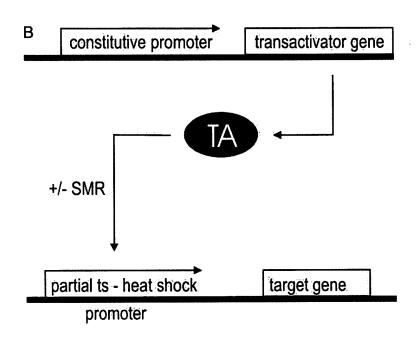
Ad - adenovirus DNA

() - gene region from which sequences can be optionally deleted

hsp70B - glvp-r promoter

- tandem or hybrid promoters





ts - transactivator - binding site SMR - small molecule regulator

TA - transactivator protein

partial ts - heat shock promoter - hybrid promoter co-activated by TA and endogenous HSF

>pShuttle 6621bp

CATCATCAATAATATACCTTATTTTGGATTGAAGCCAATATGATAATGAGGGGGGTGGAGTTTGTGACGTGG CGCGGGGCGTGGGAACGGGGGGGGGGACGTAGTAGTGTGGCGGAAGTGTGATGTTGCAAGTGTGGCGGAAC ACATGTAAGCGACGGATGTGGCAAAAGTGACGTTTTTTGGTGTGCGCCGGTGTACACAGGAAGTGACAATTT TCGCGCGGTTTTAGGCGGATGTTGTAGTAAATTTGGGCGTAACCGAGTAAGATTTGGCCATTTTCGCGGGA AAACTGAATAAGAGGAAGTGAAATCTGAATAATTTTGTGTTACTCATAGCGCGTAATACTGGTACCGCGGC CGCCTCGAGTCTAGAGATATCGAATTCAAGCTTGTCGACTCGAAGATCTGGGCGTGGTTAAGGGTGGGAAA GAATATATAAGGTGGGGGTCTTATGTAGTTTTGTATCTGTTTTTGCAGCAGCCGCCGCCGCCATGAGCACCA ACTCGTTTGATGGAAGCATTGTGAGCTCATATTTGACAACGCGCATGCCCCCATGGGCCGGGGTGCGTCAG AATGTGATGGGCTCCAGCATTGATGGTCGCCCCGTCCTGCCCGCAAACTCTACCTTGACCTACGAGAC AAGTTGACGGCTCTTTTGGCACAATTGGATTCTTTGACCCGGGAACTTAATGTCGTTTCTCAGCAGCTGTT TAGGCCCGGGACCAGCGGTCTCGGTCGTTGAGGGTCCTGTGTATTTTTTCCAGGACGTGGTAAAGGTGACT CTGGATGTTCAGATACATGGGCATAAGCCCGTCTCTGGGGTGGAGGTAGCACCACTGCAGAGCTTCATGCT GCGGGGTGGTGTTGTAGATGATCCAGTCGTAGCAGGAGCGCTGGGCGTGGTGCCTAAAAATGTCTTTCAGT AGCAAGCTGATTGCCAGGGGCAGGCCCTTGGTGTAAGTGTTTACAAAGCGGTTAAGCTGGGATGGGTGCAT ACGTGGGGATATGAGATGCATCTTGGACTGTATTTTTAGGTTGGCTATGTTCCCAGCCATATCCCTCCGGG GATTCATGTTGTGCAGAACCACCAGCACAGTGTATCCGGTGCACTTGGGAAATTTGTCATGTAGCTTAGAA GGAAATGCGTGGAAGAACTTGGAGACGCCCTTGTGACCTCCAAGATTTTCCATGCATTCGTCCATAATGAT GGCAATGGGCCCACGGGCGGCGGCCTGGGCGAAGATATTTCTGGGATCACTAACGTCATAGTTGTGTTCCA TCCGGCCCAGGGGCGTAGTTACCCTCACAGATTTGCATTTCCCACGCTTTGAGTTCAGATGGGGGGATCAT GTCTACCTGCGGGGCGATGAAGAAAACGGTTTCCGGGGTAGGGGAGATCAGCTGGGAAGAAAGCAGGTTCC TGAGCAGCTGCGACTTACCGCAGCCGGTGGGCCCGTAAATCACACCTATTACCGGGTGCAACTGGTAGTTA AGAGAGCTGCAGCTGCCGTCATCCCTGAGCAGGGGGGCCACTTCGTTAAGCATGTCCCTGACTCGCATGTT TTCCCTGACCAAATCCGCCAGAAGGCGCTCGCCGCCCAGCGATAGCAGTTCTTGCAAGGAAGCAAAGTTTT TCAACGGTTTGAGACCGTCCGCCGTAGGCATGCTTTTGAGCGTTTGACCAAGCAGTTCCAGGCGGTCCCAC AGCTCGGTCACCTGCTCTACGGCATCTCGATCCAGCATATCTCCTCGTTTCGCGGGGTTGGGGCGGCTTTCG $\tt CTGTACGGCAGTAGTCGGTCCTCGTCCAGACGGGCCAGGGTCATGTCTTTCCACGGGCGCAGGGTCCTCGT$ 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TCCGCTTCCTCGCTCACTGACTCGCTCGGTCGTTCGGCTGCGCGGCGAGCGGTATCAGCTCACAA GGCGGTAATACGGTTATCCACAGAATCAGGGGATAACGCAGGAAAGAACATGTGAGCAAAAGGCCAGCAAA AGGCCAGGAACCGTAAAAAGGCCGCGTTGCTGGCGTTTTTTCCATAGGCTCCGCCCCCTGACGAGCATCAC AAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTATAAAGATACCAGGCGTTTCCCCCTGG AAGCTCCCTCGTGCGCTCTCCTGTTCCGACCCTGCCGCTTACCGGATACCTGTCCGCCTTTCTCCCTTCGG GAAGCGTGGCGCTTTCTCATAGCTCACGCTGTAGGTATCTCAGTTCGGTGTAGGTCGTTCGCTCCAAGCTG GGCTGTGTGCACGAACCCCCGTTCAGCCCGACCGCTGCGCCTTATCCGGTAACTATCGTCTTGAGTCCAA CCCGGTAAGACACGACTTATCGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGTAG GCGGTGCTACAGAGTTCTTGAAGTGGTGGCCTAACTACGGCTACACTAGAAGGACAGTATTTGGTATCTGC TAGCGGTGGTTTTTTTTGTTTGCAAGCAGCAGATTACGCGCAGAAAAAAAGGATCTCAAGAAGATCCTTTGA TCTTTTCTACGGGGTCTGACGCTCAGTGGAACGAAAACTCACGTTAAGGGATTTTGGTCATGAGATTATCA AACTTGGTCTGACAGTTACCAATGCTTAATCAGTGAGGCACCTATCTCAGCGATCTGTCTATTTCGTTCAT CCATAGTTGCCTGACTCCCCGTCGTGTAGATAACTACGATACGGGAGGGCTTACCATCTGGCCCCAGTGCT 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$\verb|CCCGTGATATTGCTGAAGAGCTTGGCGGCGAATGGGCTGACCGCTTCCTCGTGCTTTACGGTATCGCCGCT|\\$ $\tt CCCGATTCGCAGCGCATCGCCTTCTATCGCCTTCTTGACGAGTTCTTCTGAATTTTTGTTAAAATTTTTTGTT$ AAATCAGCTCATTTTTTAACCAATAGGCCGAAATCGGCAACATCCCTTATAAATCAAAAGAATAGACCGCG ATAGGGTTGAGTGTTCCAGTTTGGAACAAGAGTCCACTATTAAAGAACGTGGACTCCAACGTCAAAGG GCGAAAAACCGTCTATCAGGGCGATGGCCCACTACGTGAACCATCACCCAAATCAAGTTTTTTTGCGGTCGA GGTGCCGTAAAGCTCTAAATCGGAACCCTAAAGGGAGCCCCCGATTTAGAGCTTGACGGGGAAAGCCGGCG AACGTGGCGAGAAAGGAAGGGAAGAAAGCGAAAGGAGCGCGCTAGGCGCCTGGCAAGTGTAGCGGTCAC GCTGCGCGTAACCACCACCCCGCGCGCTTAATGCGCCGCTACAGGGCGCGTCCATTCGCCATTCAGGATC GAATTAATTCTTAATTAA

>pGene/V5-His 7698bp

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>pXC1 9905bp

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GCATGATCGTGCTCCTGTCGTTGAGGACCCGGCTAGGCTGGCGGGGTTGCCTTACTGGTTAGCAGAATGAA TCACCGATACGCGAGCGAACGTGAAGCGACTGCTGCTGCAAAACGTCTGCGACCTGAGCAACAACATGAAT GGTCTTCGGTTTCCGTAAAGTCTGGAAACGCGGAAGTCAGCGCCCTGCACCATTATGTTCCGGA TCTGCATCGCAGGATGCTGCTGGCTACCCTGTGGAACACCTACATCTGTATTAACGAAGCGCTGGCATTGA CCCTGAGTGATTTTTCTCTGGTCCCGCCGCATCCATACCGCCAGTTGTTTACCCTCACAACGTTCCAGTAA CCGGGCATGTTCATCATCAGTAACCCGTATCGTGAGCATCCTCTCTCGTTTCATCGGTATCATTACCCCCA TGAACAGAAATCCCCCTTACACGGAGGCATCAGTGACCAAACAGGAAAAAACCGCCCTTAACATGGCCCGC TTTATCAGAAGCCAGACATTAACGCTTCTGGAGAAACTCAACGAGCTGGACGCGGATGAACAGGCAGACAT CTGTGAATCGCTTCACGACCACGCTGATGAGCTTTACCGCAGCTGCCTCGCGCGTTTCGGTGATGACGGTG AAAACCTCTGACACATGCAGCTCCCGGAGACGGTCACAGCTTGTCTGTAAGCGGATGCCGGGAGCAGACAA GCCCGTCAGGGCGCGTCAGCGGGTGTTGGCGGGTGTCGGGGCGCAGCCATGACCCAGTCACGTAGCGATAG CGGAGTGTATACTGGCTTAACTATGCGGCATCAGAGCAGATTGTACTGAGAGTGCACCATATGCGGTGTGA AATACCGCACAGATGCGTAAGGAGAAAATACCGCATCAGGCGCTCTTCCGCTTCCTCGCTCACTGACTCGC 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>pSwitch 7323bp

GACGGATCGGGAGATCATTCGAGCTTGCATGCCTGCAGGTCGAAGCGGAGTACTGTCCTCCGAGTTTAAAA GCGGAGTACTGTCCTCCGAGGATATCAGCGGAGTACTGTCCTCCGAGTCGCGAAGCGGAGTACTGTCCTCC GAGATCGATGTCGACCCCGCCCAGCGTCTTGTCATTGGCGAATTCGAACACGCAGATGCAGTCGGGGCGGC GCGGTCCGAGGTCCACTTCGCATATTAAGGTGACGCGTGTGGCCTCGAATCGCCTGGAGACGCCATCCACG CTGTTTTGACCTCCATAGAAGACACCGGGACCGATCCAGCCTCCGCGGCCGGGAACGGTGCATTGGAACGC GGATTCCCCGTGTTAATTAACAGGTAAGTGTCTTCCTCCTGTTTCCTCCTGCTGCTATTCTGCTCAACCTT CCACCAAGCTACCGGTCCACCATGGACTCCCAGCAGCCAGATCTGAAGCTACTGTCTTCTATCGAACAAGC ATGCGATATTTGCCGACTTAAAAAGCTCAAGTGCTCCAAAGAAAAACCGAAGTGCCCCAAGTGTCTGAAGA ACAACTGGGAGTGTCGCTACTCTCCCAAAACCAAAAGGTCTCCGCTGACTAGGGCACATCTGACAGAAGTG GAATCAAGGCTAGAAAGACTGGAACAGCTATTTCTACTGATTTTTCCTCGAGAAGACCTTGACATGATTTT GAAAATGGATTCTTTACAGGATATAAAAGCATTGTTAGAATTCCCGGGTGTCGACCAGAAAAAGTTCAATA AAGTCAGAGTTGTGAGAGCACTGGATGCTGTTGCTCTCCCACAGCCAGTGGGCGTTCCAAATGAAAGCCAA GCCCTAAGCCAGAGATTCACTTTTTCACCAGGTCAAGACATACAGTTGATTCCACCACTGATCAACCTGTT 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